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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: [year=2009; month=2; day=20; hr=12; min=6; sec=13; ms=887;]

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Reviewer Comments:

<210> 2

<211> 23

<212> DNA

<213> Artificial Sequence

<400> 2

gcagtcaacg aggagcgaat cag

23

For all sequences using "Artificial sequence", for numeric identifier <213>, a mandatory feature is required to explain the source of the genetic material. The feature consists of <220>, which remains blank, and <223>, which states the source of the genetic material. To explain the source, if the sequence is put together from several organisms, please list those organisms. If the sequence is made in the laboratory, please indicate that the sequence is synthesized. These errors appear in other sequences in the sequence listing. Please make all necessary changes.

Application No: 10581085 Version No: 1.0

Input Set:

Output Set:

Started: 2009-02-04 15:12:59.055
Finished: 2009-02-04 15:12:59.757
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 702 ms
Total Warnings: 4
Total Errors: 4
No. of SeqIDs Defined: 7
Actual SeqID Count: 7

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
E 224	<220>,<223> section required as <213> has Artificial sequence or Unknown in SEQID (7)

SEQUENCE LISTING

<110> DAIKIN INDUSTRIES, LTD.

<120> Method for detecting a chemical Substance with a gene-disrupted microorganism

<130> 664582

<140> 10581085

<141> 2009-02-04

<150> JP 2003-403350

<151> 2003-12-02

<160> 7

<210> 1

<211> 1287

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 1

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gatggataaa	acagacgctt	tcaaaattaa	caacacaata	gcaatcgatg	attctaaagc	180
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cgaatcaaat	ttggacaacc	tggatatggg	ttccagtaga	aggacatcgt	tggaactttt	300
ttaatataac	ctaccatagg	acacactttg	ttgttgatgt	tggaacaattc	gttaattaag	360
agtccctaaa	cggctctact	agttccaacc	tcactttggt	ttttcatttt	tttatgtttt	420
ttctagaacc	ttctttacgt	gattctcgct	cggaatccgt	caatagaatg	ttttcagtct	480
ccgtttcaat	attctgcgca	catcaatcat	tttcttacta	catacactaa	cattactcct	540
agtttaattt	aattgaattt	ttaactttct	tttcttttca	tttggcaatt	tggtctcttg	600
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aggacctcca	ttagttagag	atctgttttt	aatccattca	cctttcattc	tactctctta	1200
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gcaaacaac	aaactaaaca	aattaac				1287

<210> 2

<211> 23

<212> DNA

<213> Artificial Sequence

<400> 2

gcagtcaacg aggagcgaat cag

23

<210> 3

<211>22
 <212> DNA
 <213> Artificial Sequence

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<210> 4
 <211> 720
 <212> DNA
 <213> *Aequorea victoria*

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 aaattggaat acaactataa ctcacacaat gtatacatca tggcagacaa acaaaagaat 480
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 <212> DNA
 <213> *Saccharomyces cerevisiae*

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 tcaaaatgct cttgccactc atccagtgcg tcaatctgtt cttttgtcag atcatctaag 180
 ggatcgatag gctgatccca atctttaata acgtccagat cgaaggagtt caatgcaaga 240
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 cctttattag aatcgtttgt gtttgaggcc ccgttacctg tgagcccggg tggatcctca 480
 cttgttttaa cacctccaaa taacaagttt ttaatgaagg acatttggtc tctataatat 540
 tccgatgtac gtgtgtgtgg ctgatgagat ttagactggg tagactattt gacgcgtcta 600
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 gttctggaag catcgagcat aatacaatac aattcaacaa aaatgcgaga aggcactgat 720
 gtcttgctgt taaagaacca aaaacgcgga cactacgacc gtcttatttc cggtagaaaa 780
 agggtagata cagttgaagg aacgaagaaa attaaaatta gaaaaaaaag taaaataaaa 840
 caaggaaggt agggtaatat ggtctcgttt cctttgtcgc tccgcaaata aaggagctta 900
 ttcccgacg ctcacatggg aatttgcgcc aaatcacgga tgtggaaaac tgatcacgtg 960
 cttcgatcgc caactactga gcgctgctcc aactgatct ggcacagctt acctcgctt 1020
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tgattcttgc ccatcatatg caaaaaagta cgtacttgat atatacaaca actgtagttc	1440
agtatagcga agtttaaatt tagaag	1466

<210> 6
<211> 23
<212> DNA
<213> Artificial Sequence

<400> 6	
acgccccttc ctttttcct ttc	23

<210> 7
<211> 22
<212> DNA
<213> Artificial Sequence

<400> 7	
cttctaaatt taaacttcgc ta	22